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Synthetic 55kD i-a
48kD i-antigen pro
48 kDa immobilizat
1AG48 (G1) surface
55kD i-antigen ami
                                                                                                                                                                                                                                 (without alignments)
1837.243 Million cell updates/sec
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55 kDa immobilizat
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                                                                                                                                                                                    February 11, 2003, 19:43:19 ; Search time 33.9429 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 | SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:*
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1 MKNNILVILIISLFINQIKS......QCDFANFLSISLLLISYYLL 468
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   908470 seqs, 133250620 residues
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                                                                                                                                       sw model
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Maximum Match 100%
Listing first 45 summaries
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AAB25882
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                                                                                                                  OM protein - protein search, using
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Perfect score:
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1-antigen nfluenzee 1-antigen 1-antigen	Human EGF repeat-c Mouse laminin G1 c Novel human diagno
AAB25884 AAB25887 ABB09437 AAB25865 AAB25865 AAB10370 AAU07370 AAB19802 AAB19803 AAB19803 AAB19803 AAB19803 AAB19803 AAB19803 AAB19803 AAB19803 AAB19804 AAB19803 AAB19804 AAB19804 AAB19803	23 28 37 6
22222222222222222222222222222222222222	55
70 70 70 70 70 70 70 70 70 70 70 70 70 7	
448888888677777777777777777777777777777	0.0
373 373 23.5 20.2 20.2 20.3 5.0 20.3 5.0 198 198 198 198 188 188 188 188 188 188	178.5 175.5
11111111111111111111111111111111111111	4 4 5

55kD i-antigen protein of parasite isolate G5. ALIGNMENTS AAB25860 standard; Protein; 468 AA. 18-DEC-2000 (first entry) AAB25860; RESULT 1 AAB25860

Immobilisation antigen; i-antigen; ichthyophthirlasis; vaccine; white spot disease; freshwater fish; Immune response; infection control.

Ichthyophthirius multifiliis WO200046373-A1

04-FEB-2000; 2000WO-US02962. 10-AUG-2000

FOUND INC. (UYGE-) UNIV GEORGIA RES FOUNI (CORR) CORNELL RES FOUND INC. (CLARY) CLARK T G. (DICK/) DICKERSON H W. (LINT/) LIN T. 99US-0118634. 99US-0122372. 99US-0124905. 99US-0131121. 04-FEB-1999; 02-MAR-1999; 17-MAR-1999; 27-APR-1999;

Clark TG, Dickerson HW, Lin T;

AAY97177;

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This invention relates to movel 1-autusen priperliae sequences.

I antigens or immobilisation antigens ar common to a variety of
I antigens or immobilisation antigens ar common to a variety of
I antigens or immobilisation antigens in response to
bymenostommatid ciliates and their expression varies in response to
convironmental stimuli This invention relates to lantigens in
controlliation antigens, a protozoan which is an obligate parasite
of freshwater fish causing ichthyophthiriasis or white spot disease. The
of freshwater fish causing ichthyophthiriasis or white spot disease. The
controlling and 55 kb. Also included in the invention are
i-antigens, of 48 and 55 kb. Also included in the invention are
antibodies capable of binding to the nuclectide sequences and a method
antibodies capable of binding to the nuclectide sequences.
A composition (containing the i-antigen nuclectide of eliciting
a numure response in fish is useful for prophylaxis, treatment or for
controlling I multifiliis infection in fish. Polynuclectide or protein
vaccines comprising a portion of the amplified product encoding an
antispent i -antigen polypeptide obtained is also useful for treating or
antispent i -antigen polypeptide obtained is also useful for treating or
preventing I multifilis infection in fish. Sequences AAA97036-A97042,
and AAA97060, AAA97060 and AAA97089 represent i antigen genes and gene
(excluding AAA97060) and AAA97089 represent primers used in the
(excluding AAA97060) and AAA97089 represent primers used in the
isolation of the i-antigen gene sequences. Sequences AAB28899-B28899 and
controlling and antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFYTTKQTDWVA 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel i-antigen polypeptides and polynucleotides from Ichthyophthirlus multifillis, useful for prophylaxis and treatment of Ichthyophthirlus infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RSFTECVKCRLNEYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVA 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYV 180
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                                                                                                                                                                  This invention relates to novel i-antigen polypeptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 2540; DB 21; Length 468; Pred. No. 9.4e-194;
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                                                                                                                               Claim 3; Figure 3; 144pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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Matches 468; Conservative
              WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            468 AA;
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Gaps

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XX retrahymena thermophila expresses two major beta-tubulin genes (BTU1 and BTU2), which encode identical beta-tubulin proteins. Either of these two genes (but not both at once) can be disrupted without a detectable change con the cell phenotype. A KiSOL substitution in the BTU1 beta-tubulin in the BTU1 beta-tubulin confers increased resistance to microtubule-depolymerizing drugs and increased sensitivity to paclitaxel, a microtubule-stabilizing drug. Cells carrying the BtU1-1KiSOM allele can be transformed to paclitaxel cells carrying the BtU1-1KiSOM allele can be transformed to paclitaxel cells carrying the BtU1-1KiSOM with a wild-type BTU1 gene fragment, eliminating the need to incorporate a means for positive containing the BtU1-1KiSOM allele, BTU1: neol construct, which containing the BtU1-1KiSOM allele, BTU1: neol construct, which containing the BtU1-1KiSOM allele, BTU1: encol construct, which containing the btu1-1KiSOM allele, BTU1: encol conferring resistance to substitutes the coding region of the neol gene (conferring resistance to bushitted the patu1-1KiSOM allele, BTU1: can be used to generate BTU1 gene knockouts and for positive selection. Heterologous nucleic acids (especially and for positive selection. Heterologous nucleic acids (especially conferring resistance expression that is maintained by way of negative selection. Preferred expression that is maintained by way of negative conformation recombination mediated insertion of a heterologous nucleic acid, thereby restoring resistance to paclitaxel in the resulting crossing conferring resistance to paclitaxel in the resulting crossing conferring resistance to paclitaxel for stimulating an immune response in a vertebrate. The transgenic contains and maintained by the protein of the producing polyclonal antibodies contained by restoring resistance to paclitaxel and the resulting contains immune response in a vertebrate of the resulting contains in the presulting contains the presult of the pack packed by the packed by the packed by the packed by th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Recombinant expression systems for expressing heterologous nucleic
                                                                                                                                                     paclitaxel sensitivity; cell surface; antigen; protozoa; cillate;
live vaccine; Ichthyophthius multifillis; immobilization-antigen;
                                                                                                                                 BTU1; beta-tubulin; protein expression system; negative selection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acids and producing recombinant protein, comprises nonpathogenic protozoa such as Tetrahymena resistant to paclitaxel
                                                                                                                                                                                                                                                                                                                               /note= "Gln encoded by CAR or TAA"
                                                                                                                                                                                                    1-antigen; freshwater; fish; protozoacide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Clark TG;
                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Fig 3A; 83pp; English.
                                                                                                55 kDa immobilization antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99US-0118634
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                                                                                                                                                                                                                                                 Ichthyophthiius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaertig J, Dickerson HW,
                                                            (first entry)
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DICKERSON H W.
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N-PSDB; AAA52136.
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                                                                                                                                                                                                                                                                                                                                                                                   WO200046381-A1.
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(DICK/) DICKER
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27-APR-1999;
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                                                            04-DEC-2000
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468 AA;

Sequence

AAY97177 standard; Protein; 468 AA.

RESULT 2 AAY97177 ID AAY9

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2000-506071/45.
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Best Local S
Matches 467;
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               ö
                                                                                                                                                                                                                                                                                  en; i-antigen; ichthyophthiriasis; vaccine;
freshwater fish; immune response; infection control.
                                              NFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYV 180
                                                                                                240
                                                                                                    Gaps
                        MKNNILVILIISLFINQIKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFV 60
                               RSFTECVKCRLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVA
                                                                                                                    CPDGTISAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAGG
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               ö
    Length 468;
                                                                                                                                                                                          GIDTCTSCNKKLTSGAEANLPESAKKNIQCDFANFLSISLLLISYYLL 468
               Indels
  Score 2540; DB 21;
Pred. No. 9.4e-194;
Mismatches 0;
                                                                                                                                                                                                                                                                      Synthetic 55kD 1-antigen protein L6P
                                                                                                                                                                                                                                                                                                                                                                                       FOUND INC
                                                                                                                                                                                                                                   AAB25882 standard; Protein; 468
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                                                                                                                                                                                                                                                                                                                                                                                                                           Lin T;
 100.0%;
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CORNELL RES FOUND INC
CLARK T G.
                                                                                                                                                                                                                                                                                                                                                        99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                   Ichthyophthirius multifillis Synthetic.
                                                                                                                                                                                                                                                                                                                                             2000WO-US02962
                                                                                                                                                                                                                                                           (first entry)
      Best Local Similarity 100.
Matches 468; Conservative
                                                                                                                                                                                                                                                                                  antigen;
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                                                                                                                                                                                                                                                                                                                                                                                                      DICKERSON H W.
LIN T.
                                                                                                                                                                                                                                                                                        white spot disease;
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17-MAR-1999
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 Query Match
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                                                                                                                                                                                                                                              AAB25882;
                                                                                                                                                                                                                                                                                                                                                                                       (UYGE-)
                                                                                                                                                                                                                                                                                                                                                                                            (CORR )
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This invention relates to novel i-antigen polypeptide sequences.

I rantigens or immobilisation antigens ar common to a variety of hypothemia called ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in response to environmental stimuli. This invention relates to i-antigens in officers and inchipophthirias and inchipophthirias of i-antigens of invention includes two polypeptide and polynuclectide sequences for two i-antigens, of 48 and 55 kb. Also included in the invention are antibodies capable of binding to the nuclectide sequences and a method for identifying I. multifiliis servicypes using the nuclectide sequences.

A composition (containing the i-antigen nuclectide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifiliis infection in fish. Polynuclectide or protein vaccines comprising a portion of the amplified product encoding an antigent I rantigen polypeptide obtained is also useful for treating or preventing I multifiliis infection in fish. Sequences AAA97036-A97042, and AAA97060) and AAA97089 represent I antigen genes and gene fragments identified in the invention. Sequences AAA97036-A97046 (excluding AAA97060) and AAA97071-A97088 represent primers used in the isolation of the i-antigen gene sequences. Sequences AAB22859-B25889 and AAB2899-B25889 and AAB2899-B25889 and AAB2899-B25899 B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B25899-B2589
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Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CPDGTISAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAGG 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 2533; DB 21; Length 468;
Pred. No. 3.4e-193;
0; Mismatches 1; Indels 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GIDTCTSCNKKLTSGAEANLPESAKKNIQCDFANFLSISLLLISYYLL
                                                                                                                                                                             Example 5; Figure 14; 144pp; English.
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Similarity 99.8%;
7; Conservative (
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320 292 379

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NCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVT 176
                                                                          177 IDYVRSFTECVKCRLNFYYNGNN--GNTP----FNPG------KSQCTPCPAIKPAN 221
                   Recombinant expression systems for expressing heterologous nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     paclitaxel sensitivity, cell surface, antigen; prótozoa; ciliate;
live vaccine; Ichthyophthilus multifiliis; immobilization-antigen;
i-antigen; freshwater; fish; protozoacide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3TU1; beta-tubulin; protein expression system; negative selection;
                                                                                                                                                    222 VAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTE----CTNCAPNFYNN-----N
                                                                                                                                                                            APN---FNPG-----NSTCLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIAS
                                                                                                                                                                                                                                                    GAT-NYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCA
                                                                                                                                                                                                                                                                                                                                                                             380 LECPAGIVLIDGIISIYKQAASECVKCAANFYIIKQIDWVAGIDICISCNKKLISGAEAN
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/note= "Gln encoded by CAR, TAG or TAA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N-PSDB; AAA52134, AAA52135
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17-MAR-1999;
27-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of hymeostomatid ciliates and their expression varies in response to environmental ciliates and their expression varies in response to rentrophental ciliates and their expession varies in response to circeshwater fish causing ichthyophthiciasis or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two i-antigens, of 48 and 55 kD. Also included in the invention are invention includes two polypeptide and polynucleotide sequences for two i-antigens, of binding to the nucleotide sequences and a method of containing the i-antigen nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting a numune response in fish is useful for prophylaxis, treatment or for controlling I multifillis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an untifillis infection in fish. Sequences AAA97036-A97042, and AAA97060, AAA97065 and AAA97089 represent i-antigen gene infection in fish. Sequences AAA97036-A97086 (excluding AAA97060) and AAA97089 represent primers used in the isolation of the inventions. Sequences AAB35899-BB5889 and in the isolation of the inventions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19;
                                                                                                       Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       57 AAFVPGASTCTPCPQKKDAGAQPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECV 116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                isolation of the i-antigen gene sequences. Sequences AAB25859-B25.
AAB25893-B25906 represent i-antigen protein and peptide sequences
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) CORNELL RES FOUND INC.
) CILARK T G.
) DICKERSON H W.
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                                                                            48kD i-antigen protein sequence.
                                                                                                                                                                                                                                                                                                                    99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                      Ichthyophthirius multifiliis
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                                       (first entry)
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Matches 214; Conservative
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                                                                                                                                                                                                         WO200046373-A1
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17-MAR-1999;
27-APR-1999;
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                                       18-DEC-2000
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 AAB25859,
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(DICK/)
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Tetrahymena thermophila expresses two major beta-tubulin genes (BTU1 and BTU2), which encode identical beta-tubulin proteins. Either of these two genes (but not both at once) can be disrupted without a detectable change in the cell phenotype. A Ki350L substitution in the BTU1 beta-tubulin proteins on fers increased resistance to microtubule-depolymerizing drug. Calls carrying the BtU1-1Ki350M allele can be transformed to pacultaxel cand increased sensitivity to pacultaxel, a microtubule-stabilizing drug. Calls carrying the BtU1-1Ki350M allele can be transformed to pacultaxel cand increased sensitivity to pacultaxel a means for positive BtU1-1Ki350M allele, BTU1-1Ki350M with a wild-type BTU1 gene containing the BtU1-1Ki350M allele, BTU1: neol construct, which selection where the host organism is not a T. thermophila mutant substitutes the coding region of the neol gene (conferring resistance to parmomycin) for that of BTU1, can be used to generate BTU1 gene knockouts and for positive selection. Heterologous nucleic acids (especially concessful cell-sufface expression that is maintained by way of negative selection. Preferred expression that is maintained by way of negative selection. Preferred expression that is maintained by way of negative candid, thereby restoring resistance to pacultaxel in the resulting called proteoze are useful as live vaccines for stimulating an immune response in a vertebrate. The transgenic proteozoan host cells are also useful for producing polycloan antibodies (claimed). In particular, Tetrahymena expressing Ichthyophthius mitate effective vehicles for multifillis immobilization-antigen of freshwater fish against
acids and producing recombinant protein, comprises nonpathogenic protozoa such as Tetrahymena resistant to paclitaxel
                                                                                                                                                                            Disclosure; Fig 3A; 83pp; English.
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Immobilisation antigen; 1-antigen; lchthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.

Chthyophthirius multifiliis.

(AG48 (G1) surface protein amino acid sequence.

(first entry)

18-DEC-2000

AAB25889 standard; Protein; 409 AA.

RESULT

442 AA; Seguence

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19;
36.3%; Score 921; DB 21; Length 442;
41.8%; Pred. No. 5.8e-65;
Live 45; Mismatches 139; Indels 114; Gaps
                                                                                             57 AAFVPGASTCTPCPQKKDAGAQPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECV 116
                                                                                                                                           117 NCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVT 176
                                                                                                                                                                                                          VAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTE---CTNCAPNFYNN-----N 272
                                                                                                                                                                                                                                                                                                                                                                                                    177 TDYVRSFTECVKCRLNFYYNGNN--GNTP----FNPG------KSQCTPCPAIKPAN 221
                                                                                                                                                                                                                                                           APN---FNPG-----NSTCLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIAS 320
                                                                                                                                                                                                                                                                                                         GAT-NYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCA 379
                                                                                                                                                                                                                                                                                                                                                       380 LECPAGIVLIDGIISTYKQAASECVKCAANFYTIKQIDWVAGIDICISCNKKLISGAEAN 439
                                             1 MKNNILVILLISLFINQIKSANCPVGTETNTAGQVD----DLGTPANCVNCQKNFYYNNA 56
                                                           Conservative
             Local Similarity
                      Matches 214;
 Query Match
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Interaction regards to love I cantigen by peptides Sequences.

Interaction regards and their expression variety of hymenostomatid ciliates and their expression variety of hymenostomatid ciliates and their expression varies in response to environmental stimuli. This invention relates to 1-antigens in cirthyophthirus multifilis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthirasis or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two 1-antigens, of 48 and 55 kb. Also included in the invention are antibodies capable of binding to the nucleotide sequences and a method for identifying I. multifiliis serotypes using the nucleotide sequences. A composition (containing the 1-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifiliis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an preventing I. multifillis infection in fish. Sequences AAA97086. AAA97085 and AAA97089 represent 1-antigen genes and gene fragments identified in the invention. Sequences AAA97084.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  fragments identified in the invention. Sequences AAA97043-A97064 (excluding AAA97060) and AAA97071-A97088 represent primers used in the isolation of the 1-antigen gene sequences. Sequences AAB25859-B25889 and AAB25893-B25906 represent 1-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to novel 1-antigen polypeptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31.6%; Score 801.5; DB 21; Length 409; 39.5%; Pred. No. 1.7e-55; Live 39; Mismatches 137; Indels 111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Figure 8; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                         (UYGE-) UNIV GEORGIA RES FOUND INC.
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CLARK T G.
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99US-0122372.
99US-0124905.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DICKERSON H W.
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                                                                                                                                                                                                                  W0200046373-A1
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17-MAR-1999;
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Best Local S
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18;

23 CPVGTETNTAGQVD----DLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGAQ 78

Conservative

Matches 187;

LPESAKKNIQC---DFANFLSISLLLISYYLL 468

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Similarity

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invention relates to novel i-antigen polypeptide sequences
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02-MAR-1999;
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27-APR-1999;
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(DICK/)
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Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
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                                                                     79 PNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTCT 138
                                                           139 ACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECVKCRLNFYYNGN 198
                                                                                             199 N--GNTP----FNPG-----KSQCTPCPAIKPANVAQATLGNDATITAQCNVACPD 243
                                                                                                             121 SPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--NDSPATAGAQANLATQCSNQCPT 178
                                                                                                                                                               CLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIASGAT-NYVILQTECLNCAANFYF 341
                                                                                                                                                                                                342 DGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAAS 401
         44
                                                                                                                                                                                                           --- QGEAN
                                                                                                                              GTISAAGVNNWVAQNTE---CTNCAPNFYNN-----NAPN---FNPG------NST
                                                                                                                                                                         CVPCQINKN-DSPATAGAQANLATQCSTQCPTGTAIQDGVTLVFSNSSTQCSQCIANYFF
                                                                                                                                                                                                                                  ECVKCAANFYTTKQTDWVAGIDTCTSCNKKLTSGAEANLPESAKKNIQCDFANF 455
                                                                                                                                                                                                                                           55kD i-antigen amino acid repeat sequence SEQ ID 57.
Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                              AAB25885 standard; Peptide; 76 AA
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CLARK I G.
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99US-0122372.
99US-0124905.
99US-0131121.
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                                            45 GNOPFAAN
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02-MAR-1999;
17-MAR-1999;
27-APR-1999;
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(CLAR/)
(DICK/)
(LINT/)
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Tantigens or immobilisation antigens ar common to a variety of hymenostomatid ciliates and their expression varies in response to hymenostomatid ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens to rentification and includes two polypeptide and polymucleotide sequences. The includes two polypeptide and polymucleotide sequences. The invention includes two polypeptide and polymucleotide sequences for two i-antigens, of 48 and 55 kb. Also included in the invention are attibodies capable of binding to the nucleotide sequences and a method of for invention in containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifilial sinfection in fish. Polymucleotide or protein vaccines comprising a portion of the amplified product encoding an antigenic i-antigen polypeptide obtained is also useful for treating or preventing I. multifilial infection in fish. Sequences AAA97063, AAA97065 and AAA97069 represent i-antigen genes and gene fragments identified in the invention. Sequences AAA97043, and gene fragments identified in the invention. Sequences AAA97063, AAA97065 and AAA9701-AA97088 represent primers used in the isolation of the i-antigen gene sequences. Sequences AAB58699-B2889 and AAB58893-B25906 represent i-antigen genes and gene isolation of the i-antigen protein and peptide sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            165 CPTGTALDDGVTTDYVRSFTECVKCRLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQ 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    55kD i-antigen amino acid repeat sequence SEQ ID 55.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lin
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99US-0122372.
99US-0124905.
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CLARK T G.
DICKERSON H W.
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Best Local Similarity 100.4
Matches 76; Conservative
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This invention relates to novel i-antigen polypeptide sequences.

I antigens or immobilisation antigens ar common to a variety of hyperbear in capitates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in centromental stimuli. This invention relates to i-antigens in cof freshwater fish causing lothtyophthirias or white spot disease. The cof freshwater fish causing lothtyophthirias or white spot disease. The cof freshwater fish causing lothtyophthirias or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two antibodies capable of binding to the nucleotide sequences and a method for identifying I. multifiliis serving using the nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting a numner response in fish is useful for prophylaxis, treatment or for controlling I. multifiliis infection in fish. Polynucleotide or protein antigence i-antigen polypeptide obtained is also useful for treating or preventing I. multifiliis infection in fish. Sequences AAA97042, freaments identified in the invention. Sequences AAA97064, and AAA97060) and AAA97089 represent i-antigen gene sequences. Sequences Concluding AAA97060, and AAA97001-A97088 represent primers used in the capable of the invention and peptide sequences.
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                  multifilits, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23 CPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGAQPNPP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15.9%; Score 404; DB 21; Length 7:
100.0%; Pred. No. 7.3e-25;
Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     55kD i-antigen amino acid repeat sequence SEQ ID 58.
                                                                                   Disclosure; Figure 5b; 144pp; English.
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CORNELL RES FOUND INC.
CLARK T G.
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99US-0122372.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              83 ATANLVTQCNVK 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          72 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200046373-A1
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17-MAR-1999;
27-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB25886;
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I-antigens or immobilisation antigens are common to a variety of hymorostomatid ciliates and their expression varies in response to hymorostomatid ciliates and their expression varies in response to convironmental stimuli. This invention relates to i-antigens in ichthyophthitius multifilis, a protozoam which is an obligate parasite of freshwater fish causing ichthyophthitiasis or white spot disease. The confidence of antibodies capable of binding to the nucleotide sequences for two antibodies capable of binding to the nucleotide sequences and a method for identifying I. multifilis serotypes using the uncleotide of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I multifilis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an antigenic 1-antigen polypeptide obtained is also useful for treating or preventing I. multifilis infection in fish. Sequences AAA97060, AA97062 and AAA97060 and AAA97061 and AAA97060 and AAA97061 and AAA97060 and AAA97061 and AAA9701-A97088 represent primers used in the containing containing the containing and contained the invention. Sequences AAA97043 and gene for an impact of the invention. Sequences AAA97043 and the containing containing the containing and containing the containing and containing and containing the containing and containing the containing 
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                                                                                                                                                                     Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                         invention relates to novel 1-antigen polypeptide sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15.9%; Score 403; DB 21; Length 71; 100.0%; Pred. No. 8.6e-25; 1ve 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                55kD i-antigen amino acid repeat sequence SEQ ID 60.
                                                                                                                                                                                                                                                                       Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB25888 standard; Peptide; 72 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cchthyophthirius multifiliis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0118634.
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                                                                        Clark TG, Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best_Local Similarity 100.
Matches 71; Conservative
  (DICK/) DICKERSON H W. (LINT/) LIN T.
                                                                                                                         WPI; 2000-506071/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               301 AATLAKQCNIA 311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 AATLAKQCNIA 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      71 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200046373-A1
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02-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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10-AUG-2000
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                                                                                                                                                                                                                                                                                      δ
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                                                                                           0;
                                                                                                                                                                                                                                                                                                                                  Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                               Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                            .
0
                                                                                                                                                                                                                  Length 72;
                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                          55kD i-antigen amino acid repeat sequence SEQ ID 56.
                                                                                                                                                                                                                  15.3%; Score 389; DB 21; 1
100.0%; Pred. No. 1.1e-23;
tive 0; Mismatches 0;
                                                                                     Disclosure; Figure 5b; 144pp; English.
               (UYGE-) UNIV GEORGIA RES FOUND INC. (CORR) CORNELL RES FOUND INC. (CLAR/) CLARK T G.
                                                                                                                                                                                                                                                                                             AAB25884 standard; Peptide; 70 AA.
                                                                                                                                                                                                                                                                                                                                                  Ichthyophthirius multifiliis
  99US-0124905
99US-0131121
                                                                                                                                                                                                                                                                                                                 18-DEC-2000 (first entry)
                                                                                                                                                                                                                        Local Similarity and 72; Conservative
                                               Clark TG, Dickerson HW,
                                                                                                                                                                                                                                                           442 ESAKKNIQCDFA 453
                               DICKERSON H W.
                                                                                                                                                                                                                                                                WPI; 2000-506071/45.
                                                                                                                                                                                                          72 AA;
                                                                                                                                                                                                                                                                                                                                                             WO200046373-A1.
   17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                                                                                                                      AAB25884:
                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                    Query Match
                                     (/INIT/)
                                DICK/)
                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                    RESULT 11
                                                                                                                                                                                                                                                                                         AAB25884
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This invention relates to novel i-antigen polypeptide sequences.

I rantigens or immobilisation antigens ar common to a variety of incomposition of incomposition of incomposition of incomposition in relates to i-antigens in environmental ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in chitrophthrius multifilis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthria asis or white spot disease. The control includes two polypeptide and polynucleotide sequences for two invention includes two polypeptide and polynucleotide sequences for two antibodies capable of binding to the nucleotide sequences for two antibodies capable of binding to the nucleotide sequences and a method controlling in multifilis serotypes using the nucleotide sequences.

A composition (containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling in multifilis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an vaccines comprising a polypeptide obtained is also useful for treating or preventing I multifilis infection in fish. Sequences AAA97036-A97042, and AAA97060, AAA97065 and AAA97089 represent i-antigen genes and gene (excluding AAA97060) and AAA97089 represent primers used in the call solation of the i-antigen gene sequences. Sequences AAB28899 and AAB28893-B25906 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel i-antigen polypeptides and polynuclectides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 95 CPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNA 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 70;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14.8%; Score 375; DB 21; 100.0%; Pred. No. 1.4e-22; 1ve 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Pred. No. 1.4 ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                      (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB25887 standard; Peptide; 70 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lin T;
                                                                                                                                                                                                                                                                              CORNELL RES FOUND INC
                                                                                               99US-0122372.
99US-0124905.
99US-0131121.
04-FEB-2000; 2000WO-US02962.
                                                                 99US-0118634
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Best Local Similarity 100.0
Matches 70; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          155 ATIVAQCNVA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 ATIVAQCNVA 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      70 AA;
                                                                                                                                                                                                                                                                                                               (CLAR/) CLARK T
(DICK/) DICKERS
(LINT/) LIN T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-2000
                                                                                                      02-MAR-1999;
17-MAR-1999;
                                                                                                                                                                        27-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Clark TG,
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This invention relates to novel i-antigen polypeptide sequences.

I antigens or immobilisation antigens ar common to a variety of hymenostomatid ciliates and their expression varies in response to convince the standing standing invention relates to 1-antigens in to chityophthirius multifilis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthiriasis or white spot disease. The cinvention includes two polypeptide and polymuclectide sequences for two for indees two polypeptide and polymuclectide sequences for two for indees capable of binding to the nuclectide sequences and a method contibuting I multifilial serotypes using the nuclectide sequences.

C for identifying I multifilial serotypes using the nuclectide sequences.

C for identifying I multifilial infection in fish. Polymuclectide or protein controlling I multifilials infection in fish. Sequences AAA97086, and AAA97086, AAA97086, AAA97086, AAA97086, AAA97089 represent l antigen genes and gene cervaturing AAA97080, and AAA97089 represent l'antigen genes and gene cervaturing AAA97080) and AAA97088 represent primers used in the invention. Sequences AAA97089 and AAA97080, and AAA97080, and AAA97081 and AAA97011-A97088 represent primers used in the isolation of the invention. Sequences AAA97089-B25889 and solation of the lantigences. Sequences AAA97089 and AAA97081-A97088.
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white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel 1-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   312 CPDGTAIASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGT 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        isolation of the 1-antigen gene sequences. Sequences AAB25859-B256
AAB25893-B25906 represent 1-antigen protein and peptide sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14.7%; Score 373; DB 21;
100.0%; Pred. No. 2.1e-22;
1ve 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                                                       FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABB09437 standard; Protein; 1588 AA.
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                                                                                                                                                                                                                                                                                      CORNELL RES FOUND INC
                                                                                                                                                                             99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                   Ichthyophthirius multifillis
                                                                                                                                          04-FEB-2000; 2000WO-US02962
                                                                                                                                                                                                                                                                    (UYGE-) UNIV GEORGIA RES
                                                                                                                                                                                                                                                                                                                                                                          Clark TG, Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 100.
                                                                                                                                                                                                                                                                                                                        DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         372 ATLIAQCALE 381
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              70 AA;
                                                                       WO200046373-A1.
                                                                                                                                                                                                                                                                                                                                           LIN T.
                                                                                                                                                                             04-FEB-1999;
                                                                                                                                                                                                02-MAR-1999;
17-MAR-1999;
                                                                                                                                                                                                                                  27-APR-1999;
                                                                                                          10-AUG-2000
                                                                                                                                                                                                                                                                                                                        (DICK) (LINT/)
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                                                                                                                                                                                                                                                                                                     (CLAR/)
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The invention relates to modulating an activity of a DXR reductoisomerase enzyme of Haemophilus influenzae, comprising contacting the enzyme with a compound that modulates non-mevaloante isoprenoid blosynthesis.

"Compounds of the method is useful for treating a mammal or mammalian tissue infected with H. influenzae having DXR reductisomerase enzyme, e.g. a numan or a domestic animal. In particular, the method is useful for treating ear infections, conjunctivitis, menigitis, pneumonia, conjunctivitis, bacteraemia, sinusitis, pleural empyema, endocarditis and epiglottitis. The current sequence represents a H. influenzae DXR reductoisomerase enzyme related polypeptide sequence.

Note: The current sequence contains within it the amino acid sequence given in record ABB09436 (DXR enzyme), but this is broken up by a large insertion that appears to be accidentally inserted into the sequence.

"Consistating the DXR encoding DNR sequence represented as an amino acid
                                                                                                                                                                                                                                        241..1431
/note= "region that appears to be accidentally inserted into the sequence, consisting the DXR encoding DNA sequence represented as an amino acid sequence in three letter code"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modulating Haemophilus influenzae DXR reductoisomerase enzyme activity, useful for treating mammals or tissues infected with H. influenzae (e.g. ear infections or pneumonia) by contacting the enzyme with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           644 AGCTT-İTTAİTĞATGCCĞİAAAAACTATGGCTCGAA---ĞCTTTTAC-CAGTAĞATAG 698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                       DXR; reductoisomerase; enzyme; non-mevalonate isoprenoid; menaquinone; ubiquinone; virucide; ear infection; conjunctivitis; meningitis; pneumonia; conjunctivitis; bacteraemia; sinusitis; pleural empyema; endocarditis; epiglottitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21 ANCPVGTETNTAGQYDDLGTPANCVNCQKNFYYNNAAFVPGASTCTPCPQKKDAGAQPN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  81 PPA-----TANLVTQCNVKC------PAGTAIAGGATDYAAIITECVNCRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     699 TGAACATAATGCTATCTTTCAATCATTACCGCCAGAAGCACAAGA---AAAAATC----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8.4%; Score 213.5; DB 23; Long. Score 213.5; DB 23; Long. No. 6.2e-08; Indels 157;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yan
                                                    H. influenzae DXR related polypeptide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jaworski DD, Payne DJ, Slater-Radosti CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 40-44; 44pp; English.
                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SMIK ) SMITHKLINE BEECHAM CORP (SMIK ) SMITHKLINE BEECHAM PLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in three letter code.
                                                                                                                                                                                                                                                                                                                                                                                                                            09-AUG-2001; 2001WO-US24950
                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-AUG-2000; 2000US-223909P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25.28;
                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (e.g. ear infections or pn
modulator of its activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 25.2 hes 125; Conservative
                                                                                                                                                                                   Haemophilus influenzae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-241698/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1588 AA;
                                                                                                                                                                                                                                                                                                                                                    WO200211673-A2.
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121 NFYNENAPNFNAGAST----CTACPVNRVGGALTAGN------AATIVAQCNVACPT 167

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                                                                                                                                                                                                                                                                                                                                                           antigen; i-antigen; ichthyophthiriasis; vaccine; ease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                               993 TATTCATCCACAATCAATTATTCATCT----ATGGTACGG---TATGTTGACGCCTCAG 1045
                                                                                                                                                          954 CIGCCITITCAAT-----GCA-----AGIGCGGAAGAAAIGGAAGITAI 992
                                                                                                                      904 GTC-----GATTCAGCTACAATGATGATAAGGGCTTGGAATACATTGAGGCTCG 953
                                                                                                                                         312 CPDG-TALASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGG 370
         -----VTTDYVRSFTECVKCRLNFYYNGNNGNTPFN 205
                                                                                  903
                                                                                                    PNFYNNNAPNFNPGNSTCLPCPANKDYGAE-----ATAGGAATL----AKQCNIA 311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to novel i-antigen polypeptide sequences. I-antigens or immobilisation antigens ar common to a variety of hymenostomatid ciliates and their expression varies in response to
                                                                PGKSQCTPCPAIKPANVAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTECTNCA
                                                                                   846 CATAACACCAGAACAAGGGGTTG--CACACCCTAATTGGTCTATGGGTAAAAATTTCT
                                                                                                                                                                               371 TATLIA---QCALECPAGTVLTDGTTSTYKQAASECVKCAANFYTTKQTDWVAGIDTC--
                                                                                                                                                                                                                                                                                                                                         18kD i-antigen repeat amino acid sequence SEQ ID 12.
                                               GTTCTGGCGGACCATTCCGTTACACGCCACTTGAACAATTCACCAA-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Figure 5a; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             UNIV GEORGIA RES FOUND INC.
CORNELL RES FOUND INC.
CLARK T G.
DICKERSON H W.
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                                                                                                                                                                                                                                                                                     AAB25865 standard; Protein; 72
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                                                                                                                                                                                                                                                                                                                                                                                        Ichthyophthirius multifiliis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                              04-FEB-2000; 2000WO-US02962.
                                                                                                                                                                                                                                      1046 TCATTACTCAAATGGGA 1062
                                                                                                                                                                                                                    426 ----TSCNKKLTSGAEA 438
                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                        white spot disease;
                                                                                                                                                                                                                                                                                                                                                                                                           WO200046373-A1
                                                                                                                                                                                                                                                                                                                                                              [mmobilisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-MAR-1999;
17-MAR-1999;
27-APR-1999;
                               GTALDDG-
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controlling I multifilis a protozoan which is an obligate parasite coffreshwater fish causing ichthyophthirias; or white spot disease. The coffreshwater fish causing ichthyophthirias; or white spot disease. The convention includes two polypeptide and polynucleotide sequences for two i-antigens, of 48 and 55 kb. Also included in the invention are antibodies capable of binding to the nucleotide sequences and a method containing in multifilis serotypes using the nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I multifilis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an antigenic i-antigen polypeptide obtained is also useful for treating or preventing I multifillis infection in fish. Sequences AAA97036 AB7042, and AAA97085 and AAA97081 represent i-antigen genes and gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invention is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                             fragments identified in the invention. Sequences AAA97043-A97064 (excluding AAA97060) and AAA97071-A97088 represent primers used in the isolation of the i-antigen gene sequences. Sequences AAB28859-B25889 and AAB25893-B25906 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated nucleic acid detection reagent for detecting 1000 or more genes from Drosophila and for elucidating cell signalling and cell-cell interactions -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   382 CPAGTVLIDGITSTYKQAASECVKCAANFYTTKQIDWVAGIDTCISCNKKLISGAEANLP 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Drosophila; developmental biology; cell signalling; insecticide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 72;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8.3%; Score 212; DB 21;
53.6%; Pred. No. 1.4e-09;
tive 12; Mismatches 20;
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11-JUL-2000; 2000US-0614150
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N-PSDB; ABL08364.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PEKE ) PE CORP NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    442 ESAKKNIQC 450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 AEATQKVQC 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  72 AA;
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useful in developmental biology and in elucidating cell signalling and cell-cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA (ABBS7737-ABBS7072).

The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                          :||| || || || 1
1646 IDDLLNIANIGN------QCPPLRALKSQISRGFNCNVGEVLNMDTSD 1687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1731 TYTKEEGTKSQADCIPVCGYGTYSPTGL-----VPCLECPRNSF-----TAEPPTGG 1777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1778 FKDCQACPAQSFTYQPA-----ASNKDLCRAKCAPGTYSATGL-----APCSP 1820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1871 CVPMGHDIQCF-CPAG--FSGRRCEQDIDECASQPCYN------GGQCKDLPQGYRC- 1918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1821 CPLHHYQGAA-----GAQSCNECPSNMRTDSPASKG-----REQCKPVVCGEGACQHGGL 1870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88 VIQCNVKCPAGTAIAGGAIDYAAIITECVNCRINFYNENAPNFNAGASTCTACPVNRVGG 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              148 ALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECVKCRLNFYYNGNNGNTPFNPG 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           208 KSQCTPCPA----IKPANVAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTECTN 263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        264 CAPNEYNNNAPNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCN-1ACPDGTAIASGA 322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   323 TNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQ---CA 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 380 LECPAGTVLTDGTTSTYKQAASEC------VKC-AANFYTTKQTDWV 419
                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                      35 VDDLGTPANCVNCQKNFYINNAAAFVPGASTCTPCPQKK---DAGAQPNPPATANL---- 87
                                                                                                                                                                                                                                         Query Match
8.1%; Score 206; DB 22; Length 3396;
Best Local Similarity 22.8%; Pred. No. 6.7e-07;
Matches 103; Conservative 37; Mismatches 181; Indels 130;
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